

# SEQUENCE LISTING

<110> Meulewater, Frank  
Cornelissen, Marc  
Van Eldik, Gerben  
Jacobs, John

<120> Methods and means for delivering inhibitory RNA to plants and applications thereof

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<170> PatentIn Ver. 2.0

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<213> Artificial Sequence

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<223> Description of Artificial Sequence:cDNA copy of the nucleotide sequence of the genome of TNV-A

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<223> Description of Artificial Sequence: cDNA copy of  
the nucleotide sequence of the genome of TMV-U1

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<223> Description of Artificial Sequence: cDNA copy of  
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tcttcccgaa cgcatacgtt agtgtgacta ccgttggtcg aaacaagtaa aacaggaagg 1020  
gggttcgaat cctcctcctaa ccgcgggtaa gcggccca 1058

<210> 5

<211> 6355

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: cDNA copy of  
the nucleotide sequence of the genome of TMV-U2

<400> 5

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tggtaaaaac actctcggtt atgaccttgc aagaaggcgc atgtacgata cggccgtgga 180  
agaattttaac gcccgcgacc gtagaccaaa ggtcaacttt tccaaaacta ttagcgaaga 240  
gcaaacgctt ctagtctcca acgcgtaccc ggagttccag attacctttt ataatactca 300  
aaatgccgta cacagtttgg ctggaggttt gagagcatta gaattggaat atctgatgct 360  
acaagttccc tatggatcgc cgacatatga tataggtggg aactttgcag cacatttgtt 420

caaaggcagg gattacgtgc attgctgtat gcccaatctg gacatacgag atataatgag 480  
 gcacgaagga caaaaggact caattgagat gtatttgtcc agattgtctc gttctaacaa 540  
 ggtaattcct gagtttcaaa gggaggcttt taacagggtat gcagaagctc ccaacgaagt 600  
 ctgctgctct aaaacttttc aggattgtcg aatacatccg ccagagaata gtggtagaag 660  
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 agacgccttt gcttacaaga aaaccttggc catgttcaac actgaaagag caatctttag 1140  
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aagacttaaa attcaggggtg gctgatacca aaatcagcag tgggttggtcg tccacttaaa 6240  
 tataacgatt gtcatatctg gatccaacag ttaaaccatg tgatgggtgta tactgtggta 6300  
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<210> 6

<211> 2346

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequenc : nucleotide  
 sequence of the tomato phytoene desaturase (pds)  
 encoding cDNA

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 acaagtttcc atttaactct tcaacttcaa cccaaccaa tttatttctt taattgtgca 120  
 gaaccactcc ctatatcttc taggtgcttt cattcggtcc gaggtaagaa aagatttttg 180  
 tttctttgaa tgctttatgc cactcggtta acttctgagg tttgtggatc ttttaggcga 240  
 cttttttttt ttttgtatgt aaaatttggt tcataaatgc ttctcaacat aaatcttgac 300  
 aaagagaagg aattttacca agtathtagt ttcagaaatg gataattttc ttactgtgaa 360  
 atatccttat ggcaggtttt actggtatctt ttcagtaaaa tgcctcaaat tggacttggt 420  
 tctgctgtta acttgagagt ccaaggtagt tcagcttctc tttggagctc gaggtcgtct 480  
 tctttgggaa ctgaaagtcg agatgggtgc ttgcaaagga attcggtatg ttttgcgtgt 540  
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 aatacagtta actatttgga ggctgcattt ttatcatcaa cgttccgtgc ttctccgcgc 720  
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aagaagagtg cgagggtgaag caagtaggag aaatgttagg aaagctccta taaaaagga 2280  
tggcatgttg aagattagca tctttttaat cccaagttta aatataaagc atattttatg 2340  
gaattc 2346

<210> 7

<211> 7096

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: nucleotide  
sequence of the tobacco nitrate reductase (nia-2)  
encoding cDNA

<400> 7

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acttctattt agttttt

```

<210> 8

<211> 1839

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: nucleotide  
sequence of the tobacco nitrite reductase (nir-1)  
encoding cDNA

<400> 8

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tcaagctcca cgcaactccg ccgtctgtgg cagcgccgcc agctggtgct ccagagggtg 120  
ctgctgagag gctagaaccc agagttgagg aaaaagatgg ttattggata ctcaaggagc 180  
agtttagaaa aggcataaat cctcaagaaa aggtcaagat tgagaagcaa cctatgaagt 240  
tgttcatgga aaatggtatt gaagagcttg ctaagatacc cattgaagag atagatcagt 300  
ccaagcttac taaggatgat attgatgtta ggcttaagtg gcttggcctc ttccatagga 360  
gaaagaacca atatgggagg ttcattgatga gattgaagct tccaaatgga gtaacaacga 420  
gtgcacagac tcgatacttg gcgagtgtga taaggaaata cgggaaagaa ggatgtgctg 480  
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ggaatccagt aggaaatcct cttgctggaa ttgatccaga agaaatagta gacacagggc 660  
cttactactaa tttgctctcc caatttatca ctggcaattc acgaggcaat cccgcagttt 720  
ctaacttgcc aaggaagtgg aatccgtgcg tagtaggctc tcatgatctt tatgaacatc 780  
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taccacttgt tgtggactta ctagttaaca actttgggtg agttccacga gaaagagaag 1740  
aaacagaaga ctaataaaat ttagaatagt tgggtgatttt gctgtgttca taacatgtaa 1800  
tgtatgataa atcaatgcaa acatttctac ctacgtgag 1839

<210> 9

<211> 1294

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: cDNA of the  
beta-1,3-glucanase of *Nicotiana plumbagenifolia*

<400> 9

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cacaagttgt acaactgtac aagtcaaaaa acataagaag aatgaggctt tatgatccaa 180
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ggaatgcgat ttcttcagca gggttgcaaa acaatatcaa agtctcaagt tctgtagaca 480
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tgttgattct gcaatgataa atagaaaaaa aaaa 1294

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<210> 10

<211> 720

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: green  
fluorescent protein encoding region

<400> 10

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atggtgagca agggcgagga gctgttcacc ggggtggtgc ccactcctggt cgagctggac 60
ggcgacgtaa acggccacaa gttcagcgtg tccggcgagg gcgagggcga tgccacctac 120
ggcaagctga ccctgaagtt catctgcacc accggcaagc tgcccgtgcc ctggcccacc 180
ctcgtgacca ccctgacctc cggcgtgcag tgcttcagcc gctaccccca ccacatgaag 240
cagcagcact tcttcaagtc cgccatgccc gaaggctacg tccaggagcg caccatcttc 300
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aagctggagt acaactacaa cagccacaac gtctatatca tggccgacaa gcagaagaac 480
ggcatcaagg tgaacttcaa gatccgccac aacatcgagg acggcagcgt gcagctcgcc 540
gaccactacc agcagaacac ccccatcgcc gacggccccg tgctgctgcc cgacaaccac 600
tacctgagca cccagtccgc cctgagcaaa gaccccaacg agaagcgcga tcacatggtc 660
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<210> 11

<211> 1809

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial

Sequence: beta-glucuronidase encoding region

<400> 11

atggtccgctc ctgtagaaac cccaacccgt gaaatcaaaa aactcgacgg cctgtgggca 60  
ttcagtcctgg atcgcgaaaa ctgtggaatt gatcagcgtt ggtgggaaag cgcgttataa 120  
gaaagccggg caattgctgt gccaggcagt tttaacgata agttcgccga tgcagatatt 180  
cgtaattatg cgggcaacgt ctggtatcag cgcgaaagtct ttataccgaa aggttgggca 240  
ggccagcgta tcgtgctgcg ttctgatgag gtcactcatt acggcaaagt gtgggtcaat 300  
aatcaggaag tgatggagca tcagggcggc tatacgccat ttgaagccga tgtcacgccc 360  
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cagactatcc cgccgggaat ggtgattacc gacgaaaacg gcaagaaaaa gcagtcttac 480  
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cgcgttggcg gtaacaagaa agggatcttc actcgcgacc gcaaaccgaa gtcggcggtc 1740  
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aaacaatga 1809

<210> 12

<211> 411

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: cDNA copy of  
part of the region of a TMV-U2 variant comprising

the origin of assembly

<400> 12

ccctcgccaa ttgaactcac tgaaaaagtt gttgatgagt tcgtagatga agtaccgatg 60  
gctgtgaaac tcgaaagggt ccggaaaaca aaaaagagag tggtaggtaa taatgttaat 120  
aataagaaaa taaataatag tggtaagaag ggtttgaaag ttgaggaaat tgaggataat 180  
gtaagtgatg acgagtctat cgcgtcatcg agtacgtttt aatcaatatg ccttatacaa 240  
tcaactctcc gagccaattt gtttacttaa gttccgctta tgcagatcct gtgcagctga 300  
tcaatctgtg tacaaatgca ttaggtaacc agtttcaaac gcaacaagct aggacaacag 360  
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<210> 13

<211> 198

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: cDNA copy of  
STMV leader region

<400> 13

agtaaaactt accaatcaaa agacctaacc aacaggactg tcgtgggtcat ttatgctgtt 60  
ggggggacata gggggaaaac atattgcctt cttctacaag aggccttcag tcgccataat 120  
tacttggcgc ccaatttttg gtttcagttg ctgtttccag ctatggggag aggtaagggtt 180  
aaaccaaacc gtaaatcg 198

<210> 14

<211> 455

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: cDNA copy of  
STMV trailer region

<400> 14

gacaagtcgc cttggttatt tcgtgttggt ttaactgaac ctgcacataa gccttttggg 60  
tcgaagggtta aacgatccgc tctcgccttg agcttgaggc ggcgtatctc ttatgtcaac 120  
agagacactt tgggtctatg ttgtataaca atagatagac tcccgtttgc aagattaggg 180  
ttaacagatc ttgccgttag tctggttagc gcgtaaccgg ccttgattta tggaatagat 240  
ccattgtcca atggcctttg caatggaacg ccgacgtggc tgtataatac gtcgttgaca 300  
agtacgaaat cttgttagtg tttttccctc cacttaaatac gaagggtttt gttttggtct 360  
tcccgaacgc atacgttagt gtgactaccg ttgttcgaaa caagtaaaac aggaaggggg 420  
ttcgaatccc tccctaaccg cgggtaagcg gccca 455

<210> 15

<211> 1971

<212> DNA



<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: cDNA copy of  
part of the genome of a TMV-U1 variant, comprising  
MP and CP genes

<400> 15

ggaaacactg tgattatagc tgcattgttg gcctcgatgc ttccgatgga gaaaataatc 60  
aaaggagcct tttgtggtga cgatagtctg ctgtacttcc caaaggggtg tgagtttccg 120  
gatgtgcaac actccgcgaa tcttatgtgg aattttgaag caaaactggt taaaaaacag 180  
tatggatact tttgcggaag gtatgtaata catcacgaca gaggatgcat tgtgtattac 240  
gatcccttaa agttgatctc gaaacttggg gctaaacaca tcaaggattg ggaacacttg 300  
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aataatggtt catgagaatg agtcattgtc agaggtaaac cttctcaaag gagttaagct 660  
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